

SEQUENCE LISTING

<110> Presnell, Scott R.
Kuestner, Rolf E.
Gao, Zeren

<120> HUMAN CYTOKINE RECEPTOR

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<150> US 09/912,157

<151> 2001-07-24

<150> US 60/220,747

<151> 2000-07-26

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aartayttyg	tngayaaraa	raaytayaar	cayaarggng	ngngnmngng	nwsnggnaar	1380
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ytntgywsnc	ayytncayws	nmngnaycay	ggnytncarg	arccnggnca	rcayacnmgn	1620
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Val Phe Phe Thr Val Asn Ala Cys Leu Asn Gly Ser Gln Leu Ala Val 25
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Ala Ala Gly Gly Ser Gly Arg Ala Arg Gly Ala Asp Thr Cys Gly Trp 40
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Tyr Asp Asn Cys Thr Thr Tyr Leu Asn Pro Val Gly Lys His Val Ile 85
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Val Ala Val Thr Ile Leu Trp Ser Pro Gly Ala Leu Gly Ile Glu Phe 120
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Leu Lys Gly Phe Arg Val Ile Leu Glu Glu Leu Lys Ser Glu Gly Arg 135
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cag tgc caa caa ctg att cta aag gat ccg aag cag ctc aac agt agc 544
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Asn Leu Asn Ile Ser Gln His Gly Ser Asp Met Gln Val Ser Phe Asp	
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Thr Thr Glu Met Thr Ser Cys Leu Leu Gln Asn Val Ser Pro Gly Asp	
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Tyr Ile Ile Glu Leu Val Asp Asp Thr Asn Thr Thr Arg Lys Val Met	
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His Tyr Ala Leu Lys Pro Val His Ser Pro Trp Ala Gly Pro Ile Arg	
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Ala Val Ala Ile Thr Val Pro Leu Val Val Ile Ser Ala Phe Ala Thr	
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His Leu Asp Glu Glu Ser Ser Glu Ser Ser Thr Tyr Thr Ala Ala Leu	
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Pro Arg Glu Arg Leu Arg Pro Arg Pro Lys Val Phe Leu Cys Tyr Ser	
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650 655 660 665	

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Ser Val Pro Ser Ser Glu Leu Ser Leu Pro Leu Met Glu Gly Leu Ser	
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Arg	Pro	Lys	Val	Phe	Leu	Cys	Tyr	Ser	Ser	Lys	Asp	Gly	Gln	Asn	His
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Cys	Glu	Val	Ala	Leu	Asp	Leu	Trp	Glu	Asp	Phe	Ser	Leu	Cys	Arg	Glu
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Cys	Asn	Met	His	Gln	Phe	Ile	Asp	Glu	Glu	Pro	Asp	Trp	Phe	Glu	Lys
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 Met Ala Pro Trp Leu Gln Leu Cys Ser
 1 5

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 45 50 55

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 Phe Lys Tyr Asp Asn Cys Thr Thr Tyr Leu Asn Pro Val Gly Lys His
 60 65 70

gtg att gct gac gcc cag aat atc acc atc agc cag tat gct tgc cat 352
 Val Ile Ala Asp Ala Gln Asn Ile Thr Ile Ser Gln Tyr Ala Cys His
 75 80 85

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 Asp Gln Val Ala Val Thr Ile Leu Trp Ser Pro Gly Ala Leu Gly Ile
 90 95 100 105

gaa ttc ctg aaa gga ttt cgg gta ata ctg gag gag ctg aag tcg gag 448
 Glu Phe Leu Lys Gly Phe Arg Val Ile Leu Glu Glu Leu Lys Ser Glu
 110 115 120

gga aga cag tgc caa caa ctg att cta aag gat ccg aag cag ctc aac 496
 Gly Arg Gln Cys Gln Gln Leu Ile Leu Lys Asp Pro Lys Gln Leu Asn
 125 130 135

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caa caa gaa aat ata tat tca cat tta gat gaa gaa agc ccg gag tcg Gln Gln Glu Asn Ile Tyr Ser His Leu Asp Glu Glu Ser Pro Glu Ser 330 335 340			1123
tcc aca tac gct gcg gct ctc ccc aga gac agg ctc cgg cct cag ccc Ser Thr Tyr Ala Ala Ala Leu Pro Arg Asp Arg Leu Arg Pro Gln Pro 345 350 355			1171
aag gtc ttc ctc tgc tac tcc aat aaa gat ggc cag aat cac atg aac Lys Val Phe Leu Cys Tyr Ser Asn Lys Asp Gly Gln Asn His Met Asn 360 365 370			1219
gtg gtc cag tgt ttc gcc tat ttc ctg caa gat ttc tgt ggc tgt gag Val Val Gln Cys Phe Ala Tyr Phe Leu Gln Asp Phe Cys Gly Cys Glu 375 380 385			1267
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cac aaa gga ggc agc cgc ggc gag gcg caa ggc gag ttc ttc ctg gtg His Lys Gly Gly Ser Arg Gly Glu Ala Gln Gly Glu Phe Phe Leu Val 440 445 450			1459

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Ala	Val	Ala	Ala	Ile	Ala	Glu	Lys	Leu	Arg	Gln	Ala	Lys	Gln	Ser	Ser	
	455					460					465					
tct	gcc	gca	ctg	cgc	aag	ttc	atc	gcc	gtc	tac	ttc	gat	tat	tcc	tgt	1555
Ser	Ala	Ala	Leu	Arg	Lys	Phe	Ile	Ala	Val	Tyr	Phe	Asp	Tyr	Ser	Cys	
470					475					480					485	
gaa	ggg	gat	gta	ccc	tgc	agc	ctg	gac	ctg	agc	acc	aag	tac	aag	ctc	1603
Glu	Gly	Asp	Val	Pro	Cys	Ser	Leu	Asp	Leu	Ser	Thr	Lys	Tyr	Lys	Leu	
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Met	Asp	His	Leu	Pro	Glu	Leu	Cys	Ala	His	Leu	His	Ser	Gly	Glu	Gln	
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Glu	Val	Leu	Gly	Gln	His	Pro	Gly	His	Ser	Ser	Arg	Arg	Asn	Tyr	Phe	
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cgg	agc	aaa	tcg	ggc	cgc	tcc	ctg	tat	gtt	gcc	att	tgc	aac	atg	cac	1747
Arg	Ser	Lys	Ser	Gly	Arg	Ser	Leu	Tyr	Val	Ala	Ile	Cys	Asn	Met	His	
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Gln	Phe	Ile	Asp	Glu	Glu	Pro	Asp	Trp	Phe	Glu	Lys	Gln	Phe	Ile	Pro	
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Ser	Asp	Phe	Cys	Arg	Lys	Val	Glu	Ala	Cys	Val	Leu	Gly	Ala	Ala	Gly	
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Gln	Asp	Thr	Glu	Ala	Gln	Pro	Ser	Cys	Asp	Ser	Ala	Pro	Ala	Leu	Gln	
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ccc	ctg	tta	cac	gca	gtg	aaa	gct	ggc	agt	ccc	tca	gag	atg	cca	cgg	2083
Pro	Leu	Leu	His	Ala	Val	Lys	Ala	Gly	Ser	Pro	Ser	Glu	Met	Pro	Arg	
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gac	tca	ggc	ata	tat	gat	tct	tct	gta	ccc	tca	tca	gag	ctc	tct	ctg	2131
Asp	Ser	Gly	Ile	Tyr	Asp	Ser	Ser	Val	Pro	Ser	Ser	Glu	Leu	Ser	Leu	
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cct	ctg	atg	gag	gga	ctc	tcc	ccg	gat	cag	ata	gaa	aca	tct	tct	ctg	2179
Pro	Leu	Met	Glu	Gly	Leu	Ser	Pro	Asp	Gln	Ile	Glu	Thr	Ser	Ser	Leu	
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Thr	Glu	Ser	Val	Ser	Ser	Ser	Ser	Gly	Leu	Gly	Glu	Glu	Asp	Pro	Pro	

695	700	705	
acc ctc cct tcc aag ctc ttt gcc tct ggg gtg tcc aga gaa cat ggt			2275
Thr Leu Pro Ser Lys Leu Phe Ala Ser Gly Val Ser Arg Glu His Gly			
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tgc cac agc cac act gac gaa ctg caa gcg ctt gct cct ttg			2317
Cys His Ser His Thr Asp Glu Leu Gln Ala Leu Ala Pro Leu			
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 <213> Mus musculus

<400> 12

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Arg	Val	Ile	Leu	Glu	Glu	Leu	Lys	Ser	Glu	Gly	Arg	Gln	Cys	Gln	Gln
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Phe	Val	Lys	Ile	Val	Pro	Phe	Pro	Ser	Ile	Lys	Asn	Glu	Ser	Asn	Tyr
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His	Pro	Phe	Phe	Phe	Arg	Thr	Arg	Ala	Cys	Asp	Leu	Leu	Leu	Gln	Pro
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Asp	Asn	Leu	Ala	Cys	Lys	Pro	Phe	Trp	Lys	Pro	Arg	Asn	Leu	Asn	Ile
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Leu	Val	Asp	Asp	Ser	Asn	Thr	Thr	Arg	Lys	Ala	Ala	Gln	Tyr	Val	Val
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Lys	Ser	Val	Gln	Ser	Pro	Trp	Ala	Gly	Pro	Ile	Arg	Ala	Val	Ala	Ile
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Glu	Phe	Phe	Leu	Val	Ala	Val	Ala	Ala	Ile	Ala	Glu	Lys	Leu	Arg	Gln		
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	610					615					620						
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<211> 2217

<212> DNA

<213> Artificial Sequence

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<223> This degenerate nucleotide sequence encodes the amino acid sequence of SEQ ID NO:12.

<221> misc_feature
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